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RESULT 1
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48kD i-antigen pro
48 kDa immobilizat
1AG48 (G1) surface
55kD i-antigen ami
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55 kDa immobilizat
                                                                                                                                                                (without alignments)
1837.243 Million cell updates/sec
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an or equal to the score of the result being printed,
7 analysis of the total score distribution.
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| SIDS2/gcgdata/geneseq/geneseqp-embl/AA1980.DAT;*
| SIDS2/gcgdata/geneseq/geneseqp-embl/AA1981.DAT;*
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| SIDS2/gcgdata/geneseq/geneseqp-embl/AA1990.DAT;*
| SIDS2/gcgdata/geneseq/geneseqp-embl/AA1991.DAT;*
| SIDS2/gcdata/geneseq/geneseqp-embl/AA1991.DAT;*
                                                                                                                                February 11, 2003, 19:43:19 ; Search time 33.9429 Seconds
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GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                               Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                     908470 seqs, 133250620 residues
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                                                                                              sw model
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Maximum Match 100%
Listing first 45 summaries
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    protein search, using

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ALIGNMENTS

Immobilisation antigen; i-antigen; ichthyophthiriasis; vaccine; White spot disease; freshwater fish; Immune response; infection control. 55kD i-antigen protein of parasite isolate G5. AAB25860 standard; Protein; 468 AA (UYGE-) UNIV GEORGIA RES FOUND INC. (CCORR) CORNELL RES FOUND INC. (CLAR/) CLARK T G. (DICK/) DICKERSON H W. (LINT/) LIN T. Ichthyophthirius multifillis, 99US-0118634. 99US-0122372. 99US-0124905. 99US-0131121. 04-FEB-2000; 2000WO-US02962. 18-DEC-2000 (first entry) WO200046373-A1. 04-FEB-1999; 02-MAR-1999; 17-MAR-1999; 27-APR-1999; 10-AUG-2000 AAB25860;

g

Dickerson HW, Lin T;

Clark TG,

AAY97177;

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rnis invention relates to novel 1-autopu polypeptices sequences.

Inantigens or immobilisation antigens ar common to a variety of
I-antigens or immobilisation antigens ar common to a variety of
I-antigens of invention relates to i-antigens in
environmental stimuli. This invention relates to i-antigens in
controllative multifiliis, a protozoan which is an obligate parasite
of freshwater fish causing ichthyophthiriasis or white spot disease. The
of freshwater fish causing ichthyophthiriasis or white spot disease. The
of freshwater fish causing ichthyophthiriasis or white spot disease. The
i-antigens of 48 and 55 kb. Also included in the invention are
antibodies capable of binding to the nuclectide sequences and a method
antibodies capable of binding to the nuclectide sequences.
A composition (containing the i-antigen nuclectide or prophylaxis, treatment or for
an immune response in fish is useful for prophylaxis, treatment or for
controlling I multifiliis infection in fish. Polynuclectide or protein
vaccines comprising a portion of the amplified product encoding an
antispent i-antigen polypeptide obtained is also useful for treating or
antigent i-antigen polypeptide obtained is also useful for treating or
antigent i-antigen polypeptide obtained is also useful for treating or
antigent i-antigen polypeptide obtained is also useful for treating or
antigent and AAA97060) and AAA97089 represent trainers used in the
cerluding AAA97060) and AAA97081-A97088 represent trainers used in the
collation of the i-antigen protein and peptide sequences.
AAB25893-B25906 represent i-antigen protein and peptide sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 VQGAVATAGGTATLIAQCALECPAGTVLTDGTTSTYKQAASECVKCAANFYTTKQTDWVA 420
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel i-antigen polypeptides and polynucleotides from Ichthyophthirius multifillis, useful for prophylaxis and treatment of Ichthyophthirius infection in fish
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RSFTECVKCRLNFYYNGNNGNTPFNPGKSQCTPCPAIKPANVAQATLGNDATITAQCNVA 240
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          invention relates to novel i-antigen polypeptide sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 468;
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Pred. No. 9.4e-194;
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                                                                                                                         Claim 3; Figure 3; 144pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%;
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Matches 468; Conservative
               WPI; 2000-506071/45.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
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xx retrahymena thermophila expresses two major beta-tubulin genes (BTU1 and BTU2), which encode identical beta-tubulin proteins. Either of these two genes (but not both at once) can be disrupted without a detectable change con the cell phenotype. A KiSOL substitution in the BTU1 beta-tubulin protein confers increased resistance to microtubule-depolymerizing drugs on increased sensitivity to paclitaxel, a microtubule-depolymerizing drug. Cells carrying the BtU1-IKISOM allele can be transformed to paclitaxel cells carrying the BtU1-IKISOM allele can be transformed to paclitaxel corresistance by gene replacement of BU11-IKISOM with a wild-type BTU1 gene fragment, eliminating the need to incorporate a means for positive containing the BtU1-IKISOM allele, BTU1: neol construct, which containing the BtU1-IKISOM allele, BTU1: neol construct, which containing the btu1-IKISOM allele, BTU1: encol construct, which containing the btu1-IKISOM allele, BTU1: encol conferring resistance to substitutes the coding region of the neol gene (conferring resistance to bushit in a min gene knockouts and for positive selection. Heterologous nucleic acids (especially and for positive selection. Heterologous nucleic acids (especially conferring resistance expression that is maintained by way of negative selection. Preferred expression that is maintained by way of negative selection. Preferred expression that is maintained by way of negative condity, thereby restoring resistance to paclitaxel in the resulting resistance to paclitaxel in the resulting crossion for stimulating an immune response in a vertebrate. The transgenic contained in the protein of a producing polyclonal antibodies for stimulating an immune response in a vertebrate. The transgenic collated protein or producing polyclonal antibodies collated by I. multifiliis immobilization-antigen (i-antigen) protein on their surface are effective vehicles for vaccination of freshwater fish against
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Recombinant expression systems for expressing heterologous nucleic
                                                                                                                                                  paclitaxel sensitivity; cell surface; antigen; protozoa; cillate;
live vaccine; Ichthyophthilus multifillis; immobilization-antigen;
i-antigen; freshwater; fish; protozoacide.
                                                                                                                              BTU1; beta-tubulin; protein expression system; negative selection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        acids and producing recombinant protein, comprises nonpathogenic protozoa such as Tetrahymena resistant to paclitaxel
                                                                                                                                                                                                                                                                                                                         /note= "Gln encoded by CAR or TAA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Clark TG;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GEORGIA RES FOUND INC.
                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Fig 3A; 83pp; English.
                                                                                              55 kDa immobilization antigen.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   99US-0122372.
99US-0124905.
99US-0131121.
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                                                                                                                                                                                                                                            Ichthyophthiius multifiliis
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                                                           (first entry)
                                                                                                                                                                                                                                                                                                         Misc-difference 1..468
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DICKERSON H W.
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(GAER/) GAERTI
(DICK/) DICKER
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27-APR-1999;
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468 AA;

Sequence

AAY97177 standard; Protein; 468 AA.

RESULT 2
AAY97177
ID AAY97

3

us-09-497-967-7.rag

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2000-506071/45.
                                                                                                                                                                                      468 AA;
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Best Local Si
Matches 467;
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                                                                                                                                                                                                                                                                                               en; i-antigen; ichthyophthiriasis; vaccine;
freshwater fish; immune response; infection control.
                                                 NFYNENAPNFNAGASTCTACPVNRVGGALTAGNAATIVAQCNVACPTGTALDDGVTTDYV 180
                                                                                                 RSFTECVKCRLNFYYNGNNGNTPFNPGKSQCTPCPAIKPANVAQATLGNDATITAQCNVA 240
                                                                                                                                                                           Gaps
                         1 MKNNILVILIISLFINQIKSANCPVGTETNTAGQVDDLGTPANCVNCQKNFYYNNAAAFV 60
                                CPDGTISAAGVNNWVAQNTECTNCAPNFYNNNAPNFNPGNSTCLPCPANKDYGAEATAGG
                                                                                                                                                  AATLAKQCNIACPDGTAIASGATNYVILQTECLNCAANFYFDGNNFQAGSSRCKACPANK
                ö
    Length 468;
                                                                                                                                                                                                   GIDTCTSCNKKLTSGAEANLPESAKKNIQCDFANFLSISLLLISYYLL 468
                                                                                                                                                                                                          GIDTCTSCNKKLTSGAEANLPESAKKNIQCDFANFLSISLLISYYLL 468
                Indels
 Score 2540; DB 21;
Pred. No. 9.4e-194;
; Mismatches 0;
                                                                                                                                                                                                                                                                                  Synthetic 55kD 1-antigen protein L6P
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                                                                                                                                                                                                                                                                                                                                                                                                         FOUND INC
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 100.0%;
100.0%;
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CORNELL RES FOUND INC
CLARK T G.
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99US-0122372.
99US-0124905.
99US-0131121.
                                                                                                                                                                                                                                                                                                                 Ichthyophthirius multifillis. Synthetic.
                                                                                                                                                                                                                                                                                                                                                             2000WO-US02962
                                                                                                                                                                                                                                                                       (first entry)
             Matches 468; Conservative
                                                                                                                                                                                                                                                                                             Immobilisation antigen;
                                                                                                                                                                                                                                                                                                                                                                                                                                             Dickerson HW,
                                                                                                                                                                                                                                                                                                                                                                                                                         DICKERSON H W.
LIN T.
       Similarity
                                                                                                                                                                                                                                                                                                     white spot disease;
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02-MAR-1999;
17-MAR-1999;
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 Query Match
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This invention relates to novel i-antigen polypeptide sequences.

I-antigens or immobilisation antigens ar common to a warlety of
I-antigens or immobilisation antigens ar common to a warlety of
I-antigens or immobilisation and their expression varies in response to
environmental stimuli. This invention relates to i-antigens in
I-antigens multifilis, a protozoan which is an obligate parasite
of freshwater fish causing lichthyophthiriasis or white spot disease. The
invention includes two polypeptide and polynuclectide sequences for two
i-antigens, of 48 and 55 kb. Also included in the invention are
antibodies capable of binding to the nuclectide sequences and a method
for identifying I. multifilis sectorypes using the nuclectide sequences.
A composition (containing the i-antigen nuclectide) capable of eliciting
an immune response in fish is useful for prophylaxis, treatment or for
controlling I. multifilis infection in fish. Polynuclectide or protein
vaccines comprising a portion of the amplified product encoding an
antigent I-antigen polypeptide obtained is also useful for treating or
preventing I. multifilis infection in fish. Sequences AAA97036-A97042,
and AAA97060, AAA97060 and AAA97071-A97088 represent Primers used in the
fragments identified in the invention. Sequences AAA87060 and AAA97071-A97088 represent primers used in the
isolation of the i-antigen gene sequences.
ABESS893-BESS906 represent I-antigen gene sequences.
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Novel i-antigen polypeptides and polynucleotides from Ichthyophthirius multifiliis, useful for prophylaxis and treatment of Ichthyophthirius infection in fish
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NFYNENAPNFNAGASTCTACPVNRVGGALTAGNAATIVAQCNVACPTGTALDDGVTTDYV 180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RSFTECVKCRLNFYYNGNNGNTPFNPGKSQCTPCPAIKPANVAQATLGNDATITAQCNVA 240
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 MKNNILVILIISLFINQIKSANCPVGTETNTAGQVDDLGTPANCVNCQKNFYYNNAAAFV 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AATLAKQCNIACPDGTAIASGATNYVILQTECLNCAANFYFDGNNFQAGSSRCKACPANK
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                                                                                                                                                Example 5; Figure 14; 144pp; English
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320 292 379 439

us-09-497-967-7.rag

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NCRINFYNENAPNFNAGASTCTACPVNRVGGALTAGNAATIVAQCNVACPTGTALDDGVT 176
                   177 IDYVRSFTECVKCRLNFYYNGNN--GNTP----FNPG-----KSQCTPCPAIKPAN 221
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Recombinant expression systems for expressing heterologous nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BTU1; beta-tubulin; protein expression system; negative selection; paclitaxel sensitivity; cell surface; antigen; protozoa; ciliate; live vaccine; Ichthyophthilus multifillis; immobilization-antigen; i-antigen; freshwater; fish; protozoacide.
                                                                                                                                                    222 VAQATLGNDATITAQCNVACPDGTISAAGVNNWVAQNTE----CTNCAPNFYNN-----N
                                                                                                                                                                            APN---FNPG-----NSTCLPCPANKDYGAEATAGGAATLAKQCNIACPDGTAIAS
                                                                                                                                                                                                                                                  GAT-NYVILQTECLNCAANFYFDGNNFQAGSSRCKACPANKVQGAVATAGGTATLIAQCA
                                                                                                                                                                                                                                                                                                                                                                              380 LECPAGIVLIDGIISIYKQAASECVKCAANFYIIKQIDWVAGIDICISCNKKLISGAEAN
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/note= "Gln encoded by CAR, TAG or TAA"
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99US-0122372.
99US-0124905.
99US-0131121.
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17-MAR-1999;
27-APR-1999;
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         117
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I-antigens or immobilisation antigens ar common to a variety of hymeostomatid ciliates and their expression varies in response to environmental stimuli. This invention relates to i-antigens in rehypopthirius multifillis, a protozoan which is an obligate parasite of freshwater fish causing ichthyophthiriasis or white spot disease. The invention includes two polypeptide and polynucleotide sequences for two i-antigens, of 48 and 55 kD. Also included in the invention are invention includes two polypeptide and polynucleotide sequences and a method or i-antigens, of binding to the nucleotide sequences. A composition (containing the i-antigen nucleotide) capable of eliciting an immune response in fish is useful for prophylaxis, treatment or for controlling I. multifillis infection in fish. Polynucleotide or protein vaccines comprising a portion of the amplified product encoding an untifillis infection in fish. Sequences AAA97035-A97042, and AAA97060, AAA97065 and AAA97089 represent i-antigen gene infection in Sequences AAA97043-A97064 crangements identified in the invention. Sequences AAA97043-A97064 crandenting AAA97060) and AAA97008 represent primers used in the isolation of the inantigen sequences. Sequences AAB35899-B35889 and including AAA97060).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               19;
                                                                                                       Immobilisation antigen; i-antigen; ichthyophthiriasis; vaccine; white spot disease; freshwater fish; immune response; infection control.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel i-antigen polypeptides and polynucleotides from Ichthyophthirius multifillis, useful for prophylaxis and treatment of Ichthyophthirius infection in fish
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            57 AAFVPGASTCTPCPQKKDAGAQPNPPATANLVTQCNVKCPAGTAIAGGATDYAAIITECV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  isolation of the i-antigen gene sequences. Sequences AAB25859-B259 AAB25893-B25906 represent i-antigen protein and peptide sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels 114;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            36.3%; Score 921; DB 21; Length 442;
41.8%; Pred. No. 5.8e-65;
.ive 45; Mismatches 139; Indels 11.
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                                                                                                                                                                                                                                                                                                                                                                                                                 FOUND INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 1; Figure 1; 144pp; English
                                                                            48kD i-antigen protein sequence.
                                                                                                                                                                                                                                                                                                                                                                                                             (UYGE-) UNIV GEORGIA RES FOUNI
(CORR ) CORNELL RES FOUND INC.
(CLAR/) CLARK T G.
                                                                                                                                                                                                                                                                                                                     99US-0118634.
99US-0122372.
99US-0124905.
99US-0131121.
                                                                                                                                                                      Ichthyophthirius multifiliis
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                                       (first entry)
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Matches 214; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Dickerson HW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DICKERSON H W.
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17-MAR-1999;
27-APR-1999;
                                       18-DEC-2000
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 AAB25859
                                                                                                                                                                                                                                                                                                                                                                                                                                                  (CLAR/)
(DICK/)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (LINT/)
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Tetrahymena thermophila expresses two major beta-tubulin genes (BTU1 and BTU2), which encode identical beta-tubulin proteins. Either of these two genes (but not both at once) can be disrupted without a detectable change in the cell phenotype. A Ki350L substitution in the BTU1 beta-tubulin protein confers increased resistance to microtubule-depolymerizing drugs and increased sensitivity to paclitate, a microtubule-stabilizing drug. Cells carrying the Btu1-1Ki350M allele can be transformed to paclitaxel crasstance by gene replacement of Btu1-1Ki350M with a wild-type BTU1 gene for fragment, eliminating the need to incorporate a means for poslitave containing the Btu1-1Ki360M allele, BTU1: neol construct, which substitutes the host organism is not a T. thermophila mutant containing the Btu1-1Ki360M allele, BTU1: neol construct, which substitutes the coding region of the neol gene (conferring resistance to parmomycin) for that of BTU1, can be used to generate BTU1 gene for parmomycin) for that of BTU1, can be used to generate BTU1 gene for concessful cell-sufface expression that is maintained by way of negative selection. Preferred expression that is maintained by way of negative selection. Preferred expression that is maintained by way of negative concessful cell-sufface expression that is maintained by way of negative the monologous recombination-mediated insertion of a heterologous nucleic acid, thereby restoring resistance to paclitaxel in the resulting concession host. Transgenic cliated proteczea are useful as live vaccines for stimulating an immune response in a vertebrate. The transgenic concerns are also useful for producing polyclonal antibodies cliated to protect the concession host cells are also useful for producing polyclonal antibodies cliated to protect the concession host cells are also useful for producing polyclonal antibodies cliated to the concession host cells are also useful for producing and the concession protects and particular the producing polyclonal antibodies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (claimed). In particular, Terrahymena expressing Ichthyophthilus multifillis immobilization-antigen (1-antigen) protein on their surface are effective vehicles for vaccination of freshwater fish against
acids and producing recombinant protein, comprises nonpathogenic protozoa such as Tetrahymena resistant to paclitaxel
                                                                                                                                                                                                                           Disclosure; Fig 3A; 83pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  infection by I. multifilits.
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442 AA; Seguence

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19;
                                                                                     57 AAFVPGASTCTPCPQKKDAGAQPNPPATANLVTQCNVKCPAGTAIAGGATDYAAIITECV 116
                                                                                                                                117 NCRINFYNENAPNFNAGASTCTACPVNRVGGALTAGNAATIVAQCNVACPTGTALDDGVT 176
                                                                                                                                                                                          ; Score 921; DB 21; Length 442;
; Pred. No. 5.8e-65;
45; Mismatches 139; Indels 114; Gaps
                                                                                                                                               177 TDYVRSFTECVKCRLNFYYNGNN--GNTP----FNPG------KSQCTPCPAIKPAN 221
                                                                                                                                                                                                                        VAQATLGNDATITAQCNVACPDGTISAAGVNNWVAQNTE---CTNCAPNFYNN-----N 272
                                                                                                                                                                                                                                      APN---FNPG-----NSTCLPCPANKDYGAEATAGGAATLAKQCNIACPDGTAIAS 320
                                                                                                                                                                                                                                                                                 GAT-NYVILQTECLNCAANFYFDGNNFQAGSSRCKACPANKVQGAVATAGGTATLIAQCA 379
                                                                                                                                                                                                                                                                                                                              380 LECPAGIVLIDGIISTYKQAASECVKCAANFYTIKQIDWVAGIDICISCNKKLISGAEAN 439
                                         1 MKNNILVILIISLFINQIKSANCPVGTETNTAGQVD----DLGTPANCVNCQKNFYYNNA 56
                                                     LPESAKKNIQC --- DFANFLSISLLLISYYLL 468
                                                                                                                                                                                                                                                                                                                                                                                                                     36.38;
           41.8%;
                    Conservative
           Similarity
                   Matches 214;
Query Match
           Local
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18;

23 CPVGTETNTAGQVD----DLGTPANCVNCQKNFYYNNAAAFVPGASTCTPCPQKKDAGAQ 78

31.6%; Score 801.5; DB 21; Length 409; 39.5%; Pred. No. 1.7e-55; Live 39; Mismatches 137; Indels 111;

Conservative

Matches 187;

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Query Match Best Local S

Similarity

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This invention relates to novel i-antigen polypeptide sequences.

I-antigens or immobilisation antigens ar common to a variety of hypothemics to immobilisation antigens ar common to a variety of the common control of immobilisation and their expression varies in response to environmental stimuli. This invention relates to i-antigens in Inthyophthirius multifilis, a protozoan which is an obligate parasite of freshwater fish causing ichthyophthirias or white spot disease. The invention includes two polypeptide and polynuclectide sequences for two i-antigens, of 48 and 55 kb. Also included in the invention are antipodies capable of binding to the nuclectide sequences and a method for identifying I. multifillis susciples using the nuclectide sequences. A composition (containing the i-antigen nuclectide) capable of allciting an immune response in fish is useful for prophylaxis, treatment or for controlling I. multifillis infection in fish. Polynuclectide or protein vaccines comprising a portion of the amplified product encoding an antigenic i-antigen polypeptide obtained is also useful for treating or preventing I. multifillis infection in fish. Sequences AAA97060, AAA97065 and AAA97060 represent I antigen genes and gene in frame and managing and persent antigen genes and gene in antigen genes and gene in antigen genes and gene in antigen genes and sequences and AAA97060.
                                                                                                                                                         Immobilisation antigen; 1-antigen; 1chthyophthiriasis; vaccine;
white spot disease; freshwater fish; immune response; infection control.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      fragments identified in the invention. Sequences AAA77043-A97064 (excluding AAA97060) and AAA97071-A97088 represent primers used in the isolation of the 1-antigen gene sequences. Sequences AAB25859-B25889 and AAB25893-B25806 represent 1-antigen protein and peptide sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel i-antigen polypeptides and polynucleotides from Ichthyophthirius multifiliis, useful for prophylaxis and treatment of Ichthyophthirius infection in fish
                                                                                                                   IAG48 (G1) surface protein amino acid sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Figure 8; 144pp; English.
AAB25889 standard; Protein; 409 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  UYGE-) UNIV GEORGIA RES FOUND INC.
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                                                                                                                                                                                                                    Chthyophthirius multifiliis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORNELL RES FOUND INC
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99US-0122372.
99US-0124905.
99US-0131121.
                                                                                                                                                                                                                                                                                                                                     04-FEB-2000; 2000WO-US02962.
                                                                             (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Dickerson HW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DICKERSON H W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2000-506071/45.
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                                                                                                                                                                                                                                                        40200046373-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LIN T.
                                                                           18-DEC-2000
                                                                                                                                                                                                                                                                                                                                                                                04-FEB-1999;
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(CLAR/)
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WPI; 2000-506071/45.
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(DICK/)
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Immobilisation antigen; i-antigen; ichthyophthirlasis; vaccine; white spot disease; freshwater fish; immune response; infection control.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel i-antigen polypeptides and polynucleotides from Ichthyophthirius multifillis, useful for prophylaxis and treatment of Ichthyophthirius infection in fish
                                                                PNPPATANLVTQCNVKCPAGTAIAGGATDYAAIITECVNCRINFYNENAPNFNAGASTCT 138
                                                       139 ACPVNRVGGALTAGNAATIVAQCNVACPTGTALDDGVTTDYVRSFTECVKCRLNFYYNGN 198
                                                                                       199 N--GNTP----FNPG-----KSQCTPCPAIKPANVAQATLGNDATITAQCNVACPD 243
                                                                                                     121 SPQGEAPGVQVFAAGAAAAGVAAVTSQCVPCQLNK--NDSPATAGAQANLATQCSNQCPT 178
                                                                                                                                                      CLPCPANKDYGAEATAGGAATLAKQCNIACPDGTAIASGAT-NYVILQTECLNCAANFYF 341
                                                                                                                                                                                    342 DGNNFQAGSSRCKACPANKVQGAVATAGGTATLIAQCALECPAGTVLTDGTTSTYKQAAS 401
        44
                                                                                                                                                                                               --- QGEAN
                                                                                                                      GTISAAGVNNWVAQNTE---CTNCAPNFYNN-----NAPN---FNPG------NST
                                                                                                                                                              CVPCQINKN-DSPATAGAQANLATQCSTQCPTGTAIQDGVTLVFSNSSTQCSQCIANYFF
                                                                                                                                                                                                                   ECVKCAANFYTTKQTDWVAGIDTCTSCNKKLTSGAEANLPESAKKNIQCDFANF 455
                                                                                                                                                                                                                            55kD i-antigen amino acid repeat sequence SEQ ID 57.
Disclosure; Figure 5b; 144pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (UYGE-) UNIV GEORGIA RES FOUND INC.
                                                                                                                                                                                                                                                                            AAB25885 standard; Peptide; 76 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORNELL RES FOUND INC
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99US-0122372.
99US-0124905.
99US-0131121.
                                                                                                                                                                                                                                                                                                                                                                  Ichthyophthirius multifiliis.
                                                                                                                                                                                                                                                                                                                                                                                                                  2000WO-US02962.
                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          HW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DICKERSON H W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Dickerson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2000-506071/45.
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                                         45 GNOPFAAN-
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02-MAR-1999;
17-MAR-1999;
27-APR-1999;
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(CLAR/)
(DICK/)
(LINT/)
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1 Introduction tenders to mover a margin proper sequences.

I manipulates and their expression varies in response to hymenostomatid ciliates and their expression varies in response to environmental stimuli. This invention relates to i-antigens in response to environmental stimuli. This invention includes the protozoan which is an obligate parasite of Enthyophthirius multifilis; a protozoan which is an obligate parasite of inchides two polypeptide and polymucleotide sequences. The invention includes two polypeptide and polymucleotide sequences for two i-antigens, of 48 and 55 kb. Also included in the invention are antibodies capable of binding to the nucleotide sequences and a method of for invention in containing the i-antigen nucleotide) capable of eliciting an immune response in fish is useful for prophylaxis, treatment or for controlling I. multifilisis infection in fish. Polymucleotide or protein vaccines comprising a portion of the amplified product encoding an antigenic i-antigen polypeptide obtained is also useful for treating or preventing I. multifilisis infection in fish. Sequences AAA97063, AAA97065 and AAA97089 represent i-antigen genes and gene fragments identified in the invention. Sequences AAA97035-A97064 and AAA97060, AAA97071-A97088 represent primers used in the isolation of the i-antigen gene sequences. Sequences AAB58899 and isolation of the i-antigen gene and peptide sequences.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            165 CPTGTALDDGVTTDYVRSFTECVKCRLNFYYNGNNGNTPFNPGKSQCTPCPAIKPANVAQ 224
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 CPTGTALDDGVTTDYVRSFTECVKCRLNFYNGNNGNTPFNPGKSQCTPCPALKPANVAQ 60
invention relates to novel i-antigen polypeptide sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       16.7%; Score 424; DB 21; Length 76; 100.0%; Pred. No. 2e-26; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         55kD i-antigen amino acid repeat sequence SEQ ID 55.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAB25883 standard; Peptide; 72 AA.
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99US-0122372.
99US-0124905.
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DICKERSON H W.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       76 AA;
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27-APR-1999;
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02-MAR-1999;
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This invention relates to novel i-antigen polypeptide sequences.

I antigens or immobilisation antigens ar common to a variety of hymoroscomatid cilitates and their expression varies in response to environmental stimuli. This invention relates to i-antigens in relationship and their caps and their sequences to institute the confirmation includes two polypeptide and polynucleotide sequences for two invention includes two polypeptide and polynucleotide sequences for two antibodies capable of binding to the nucleotide sequences and a method for identifying I. multifiliis servitypes using the nucleotide sequences.

C and animume response in fish is useful for prophylaxis, treatment or for controlling I. multifiliis infection in fish. Polynucleotide or protein antigent i-antigen polypeptide obtained is also useful for treating or antigent i-antigen polypeptide obtained is also useful for treating or preventing I. multifiliis infection in fish. Sequences AAA97042, fragments identified in the invention. Sequences AAA97063, and AAA97063 and AAA97089 represent I antigen genes and gene captualion of the invention. Sequences AAB28899-BS889 and the i-antigen gene sequences. Sequences AAB28899-BS889 and the control of the invention and peptide sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Immobilisation antigen; i-antigen; ichthyophthiriasis; vaccine;
White spot disease; freshwater fish; immune response; infection control.
                 multifillis, useful for prophylaxis and treatment of Ichthyophthirius infection in fish -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          23 CPVGTETNTAGQVDDLGTPANCVNCQKNFYYNNAAAFVPGASTCTPCPQKKDAGAQPNPP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15.9%; Score 404; DB 21; Length 7 100.0%; Pred. No. 7.3e-25; Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        55kD 1-antigen amino acid repeat sequence SEQ ID 58.
                                                                                 Disclosure; Figure 5b; 144pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAB25886 standard; Peptide; 71 AA.
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99US-0122372.
99US-0124905.
99US-0131121.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ichthyophthirius multifiliis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            04-FEB-2000; 2000WO-US02962
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity 100.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           61 ATANLVTOCNVK 72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        83 ATANLVTQCNVK 94
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         72 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WO200046373-A1
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17-MAR-1999;
27-APR-1999;
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Matches
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(UYGE-) UNIV GEORGIA RES FOUND INC. (CORR) CORNELL RES FOUND INC. (CLAR/) CLARK T G.

I-antigens or immobilisation antigens are common to a variety of hymorostomatid ciliates and their expression varies in response to hymorostomatid ciliates and their expression varies in response to convironmental stimuli. This invention relates to i-antigens in ichthyophthitius multifilis, a protozoan which is an obligate parasite of freshwater fish causing ichthyophthitiasis or white spot disease. The confraint includes two polypeptide and polymorecetide sequences for two invention includes two polypeptide and polymorecetide sequences for two antigens, of 48 and 55 kb. Also included in the invention are method for identifying I. multifilis serotypes using the uncleotide sequences. A composition (containing the i-antigen nucleotide) capable of eliciting an immune response in fish is useful for prophylaxis, treatment or for controlling I. multifilis infection in fish. Polymorleotide or protein vaccines comprising a portion of the amplified product encoding an antigenic 1-antigen polypeptide obtained is also useful for treating or preventing I. multifilis in facction in fish. Sequences AAA97063-A97042, and AAA97060, AAA97065 and AAA97089 represent 1-antigen genes and gene fragments identified in the invention. Sequences AAA97043-A97064 and AAA97060, AAA97065 and AAA97081-A97088 represent primers used in the collation of the invention. Sequences AAA97043-A97064 and collation of the invention. Sequences AAA97043-A97064 and collation of the invention. Sequences AAA97043-A97064 in the collation of the invention of the collation of the invention of the collation of the invention. (excluding AAA97060) and AAA97071-A97088 represent primers used in the isolation of the 1-antigen gene sequences. Sequences AAB25859-B25889 and AAB25893-B25906 represent 1-antigen protein and peptide sequences. Immobilisation antigen; i-antigen; ichthyophthiriasis; vaccine; White spot disease; freshwater fish; Immune response; infection control. Novel i-antigen polypeptides and polynucleotides from Ichthyophthirius multifillis, useful for prophylaxis and treatment of Ichthyophthirius infection in fish -Gaps This invention relates to novel 1-antigen polypeptide sequences ö 15.9%; Score 403; DB 21; Length 71; 100.0%; Pred. No. 8.6e-25; 1ve 0; Mismatches 0; Indels 55kD i-antigen amino acid repeat sequence SEQ ID 60. Disclosure; Figure 5b; 144pp; English. AAB25888 standard; Peptide; 72 AA. Ë Lin chthyophthirius multifiliis 99US-0118634. 04-FEB-2000; 2000WO-US02962 (first entry) Query Match
Best Local Similarity 100.
Matches 71; Conservative Clark TG, Dickerson HW, (DICK/) DICKERSON H W. (LINT/) LIN T. WPI; 2000-506071/45 301 AATLAKQCNIA 311 61 AATLAKOCNIA 71 71 AA; WO200046373-A1 .8-DEC-2000 04-FEB-1999; 02-MAR-1999; 10-AUG-2000 Sequence AAB25888; RESULT 10 AAB25888 ò g ŏ g ö ö Length 72;

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10-AUG-2000
                                                CORR
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                                                                                                                                                                                                                                                                                                                              Immobilisation antigen; i-antigen; ichthyophthiriasis; vaccine; white spot disease; freshwater fish; immune response; infection control.
                                                              Novel i-antigen polypeptides and polynucleotides from Ichthyophthirius multifillis, useful for prophylaxis and treatment of Ichthyophthirius infection in fish
                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                         ..
                                                                                                                                                                                                              Length 72;
                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                     55kD i-antigen amino acid repeat sequence SEQ ID 56.
                                                                                                                                                                                                              15.3%; Score 389; DB 21; 1
illarity 100.0%; Pred. No. 1.1e-23;
Conservative 0; Mismatches 0;
                                                                                   Disclosure; Figure 5b; 144pp; English
               (UYGE-) UNIV GEORGIA RES FOUND INC. (CORR ) CORNELL RES FOUND INC. (CLAR/) CLARK T G.
                                                                                                                                                                                                                                                                                         AAB25884 standard; Peptide; 70 AA.
                                                                                                                                                                                                                                                                                                                                             Ichthyophthirius multifiliis
  99US-0124905.
                                                                                                                                                                                                                                                                                                             18-DEC-2000 (first entry)
                                               Clark TG, Dickerson HW,
                                                                                                                                                                                                                                                       442 ESAKKNIQCDFA 453
                               DICKERSON H W.
                                                                                                                                                                                                                                                             61 ESAKKNIQCDFA 72
                                                      WPI; 2000-506071/45.
                                                                                                                                                                                                                     Similarity
                                                                                                                                                                                                       72 AA;
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  17-MAR-1999;
27-APR-1999;
                                                                                                                                                                                                                                                                                                   AAB25884;
                                                                                                                                                                                                        Sequence
                                                                                                                                                                                                                 Query Match
                                     LINT/)
                          (CLAR/)
                                DICK/)
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This invention relates to novel i-antigen polypeptide sequences.

I antigens or immobilisation antigens ar common to a variety of incomposition of incomposition of incomposition of incomposition in relates to i-antigens in environmental ciliates and their expression varies in response to environmental stimuli. This invention relates to i-antigens in chithyophthirius multifilis, a protozoan which is an obligate parasite of freshwater fish causing ichthyophthiriasis or white spot disease. The control includes two polypeptide and polynucleotide sequences for two invention includes two polypeptide and polynucleotide sequences for two introduces, of 48 and 55 kb. Also included in the invention are antibodies capable of binding to the nucleotide sequences and a method animume response in fish is useful for prophylaxis, treatment or for an immune response in fish is useful for prophylaxis, treatment or for controlling I multifilis infection in fish. Polynucleotide or protein vaccines comprising a portion of the amplified product encoding an vaccines comprising a portion of the amplified product encoding an vaccines comprising a polypeptide obtained is also useful for treating or preventing I multifillis infection in fish. Sequences AAA97036-A97042, and AAA97060, AAA97065 and AAA97089 represent i-antigen gene sequences and gene (excluding AAA97060) and AAA97089 represent primers used in the call of the i-antigen gene sequences. Sequences AAB25859-B25809 and AAB25893-B25906 represent i-antigen protein and peptide sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel i-antigen polypeptides and polynuclectides from Ichthyophthirius multifillis, useful for prophylaxis and treatment of Ichthyophthirius infection in fish
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Figure 5b; 144pp; English.
                                                                                                                                                                                                                                  (UYGE-) UNIV GEORGIA RES FOUND INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                     Lin T;
                                                                                                                                                                                                                                                                      CORNELL RES FOUND INC
                                                                                               99US-0122372.
99US-0124905.
99US-0131121.
04-FEB-2000; 2000WO-US02962.
                                                               99US-0118634
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Best Local Similarity luv..
To 70; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                            Dickerson HW,
                                                                                                                                                                                                                                                                                                                                      DICKERSON H W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2000-506071/45.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              155 ATIVAQCNVA 164
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 61 ATIVAQCNVA 70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        70 AA;
                                                                                                                                                                                                                                                                                                      (CLAR/) CLARK T
(DICK/) DICKERS
(LINT/) LIN T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           18-DEC-2000
                                                                                            02-MAR-1999;
17-MAR-1999;
27-APR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                            Clark TG,
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This invention relates to novel i-antigen polypeptide sequences.

Tantigens or immobilisation antigens ar common to a variety of i-antigens or immobilisation antigens ar common to a variety of hymenostomatid ciliates and their expression varies in response to control stimular studii. This invention relates to i-antigens in chithyophthiria stimular is an unitifilis, a motocoan which is an obligate parasite of freshwater fish causing ichthyophthiriasis or white spot disease. The cinvention includes two polypeptide and polymuclectide sequences for two family or multifilis services in the invention are antigens, of 48 and 55 kb. Also included in the invention are controlling. In willifilitis services using the nuclectide sequences.

Composition (containing the i-antigen nuclectide) capable of eliciting an immune response in fish is useful for prophylaxis, treatment or for controlling I multifilis infection in fish. Polymuclectide or protein controlling I multifilis infection in fish. Sequences AAA97066, AAA97066, AAA97066, AAA97066, AAA97066, AAA97066, AAA97060, AAA97060, and AAA97069 represent l-antigen genes and gene cerval and AAA97060 and AAA97071-A97088 represent primers used in the invention. Sequences AAA97064 in the invention. Sequences AAA97064 in the invention of certain primers used in the isolation of the invention. Sequences AAA97069 and AAA9701-A97088 represent primers used in the isolation of the invention.
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white spot disease; freshwater fish; immune response; infection control.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel 1-antigen polypeptides and polynucleotides from Ichthyophthirius multifiliis, useful for prophylaxis and treatment of Ichthyophthirius infection in fish -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        312 CPDGTAIASGATNYVILQTECLNCAANFYFDGNNFQAGSSRCKACPANKVQGAVATAGGT 371
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               isolation of the 1-antigen gene sequences. Sequences AAB25859-B256
AAB25893-B25906 represent 1-antigen protein and peptide sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       14.7%; Score 373; DB 21;
100.0%; Pred. No. 2.1e-22;
11ve 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Figure 5b; 144pp; English.
                                                                                                                                                                                                                                                                         (UYGE-) UNIV GEORGIA RES FOUND INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABB09437 standard; Protein; 1588 AA.
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                                                                                                                                                                                                                                                                                           CORNELL RES FOUND INC
CLARK T G.
                                                                                                                                                                                 99US-0118634.
99US-0122372.
99US-0124905.
99US-0131121.
                                    Ichthyophthirius multifiliis
                                                                                                                                             04-FEB-2000; 2000WO-US02962
                                                                                                                                                                                                                                                                                                                                                                                   Clark TG, Dickerson HW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity 100.
                                                                                                                                                                                                                                                                                                                               DICKERSON H W.
                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2000-506071/45.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 372 ATLIAQCALE 381
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                                                                        WO200046373-A1.
                                                                                                                                                                                                                                                                                                                                                  LIN T.
                                                                                                                                                                                                                                                                                                                CLARK
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17-MAR-1999;
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The invention relates to modulating an activity of a DXR reductoisomerase enzyme of Haemophilus influenzae, comprising contacting the enzyme with a compound that modulates non-mevalonate isopremoid blosynthesis.

Compound that modulates non-mevalonate isopremoid blosynthesis.

Synthesis of menaquinone or ubiquinone. Compounds of the invention act as virucides. The method is useful for treating a mammal or mammalian tissue infected with H. influenze having DXR reductoisomerase enzyme, e.g. a neating ear infections, conjunctivitis, meningitis, pneumonia, conjunctivitis, bacteraemia, sinusitis, pleural empyema, endocarditis and reductoisomerase enzyme related polypeptide sequence.

Note: The current sequence contains within it the amino acid sequence given in record ABBOQ436 (DXR enzyme), but this is broken up by a large insertion that appears to be accidentally inserted into the sequence, consisting the DXR encoding DNA sequence represented as an amino acid sequence in three letter code.
                                                                                                                                                                                                                                               241..1431
/note= "region that appears to be accidentally inserted into the sequence, consisting the DXR encoding DNA sequence represented as an amino acid sequence in three letter code"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Modulating Haemophilus influenzae DXR reductoisomerase enzyme activity, useful for treating mammals or tissues infected with H. influenzae (e.g. ear infections or pneumonia) by contacting the enzyme with a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               644 AGCTT-İTTAİTĞATGCCĞİAAAAACTATGGCTCGAA---ĞCTTTTAC-CAGTAĞATAG 698
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             21 ANCPVGTETNTAGQVDDLGTPANCVNCQKNFYYNNAAAFVPGASTCTPCPQKKDAGAQPN 80
                                                                                          DXR; reductoisomerase; enzyme; non-mevalonate isoprenoid; menaquinone; ubiquinone; virucide; ear infection; conjunctivitis; meningitis; pneumonia; conjunctivitis; bacteraemia; sinusitis; pleural empyema; endocarditis; epiglottitis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        81 PPA-----TANLVTQCNVKC------PAGTAIAGGATDYAAIITECVNCRI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            699 TGAACATAATGCTATCTTTCAATCATTACCGCCAGAAGCACAAGA---AAAAATC----
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           205; Indels 157;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 23; Length 1588;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Yan
                                                      H. influenzae DXR related polypeptide sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                8.4%; Score 213.5; DB 2
25.2%; Pred. No. 6.2e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Jaworski DD, Payne DJ, Slater-Radosti CE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 40-44; 44pp; English.
                                                                                                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (SMIK ) SMITHKLINE BEECHAM CORP (SMIK ) SMITHKLINE BEECHAM PLC.
                                                                                                                                                                                                                                                                                                                                                                                                                                         09-AUG-2001; 2001WO-US24950
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             09-AUG-2000; 2000US-223909P
                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (e.g. ear infections or pn
modulator of its activity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity 25.2
Matches 125; Conservative
                                                                                                                                                                                         Haemophilus influenzae.
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                                                                                                                                                                                                                                                                                                                                                              WO200211673-A2.
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121 NFYNENAPNFNAGAST----CTACPVNRVGGALTAGN------AATIVAQCNVACPT 167

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Best Loca
Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               antigen; i-antigen; ichthyophthiriasis; vaccine; ease; freshwater fish; immune response; infection control.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Novel i-antigen polypeptides and polynucleotides from Ichthyophthirius multifiliis, useful for prophylaxis and treatment of Ichthyophthirius infection in fish
                                                                                                                                                                                                                                                                                                                                                                                                                                     993 TATTCATCCACAATCAATTATTCATTCT----ATGGTACGG---TATGTTGACGCCTCAG 1045
                                                                                                                                                                                                                                                                                                              312 CPDG-TAIASGATNYVILQTECLNCAANFYFDGNNFQAGSSRCKACPANKVQGAVATAGG 370
                                                                                                                                                                                                                                                                                                                                                     954 CIGGCITITCAAT-----GCA-----AGIGCGGAAGAAAIGGAAGITAI 992
                                                               -----VTTDYVRSFTECVKCRLNFYYNGNNGNTPFN 205
                                                                                                                                                                                                                             PNFYNNNAPNFNPGNSTCLPCPANKDYGAE-----ATAGGAATL----AKQCNIA 311
                                                                                                                                                                                                                                                                    904 GIC------GATICAGCIACAAIGAIGAAIAAGGGCIIGGAAIACAIIGAGGCICG 953
                     -----GGTTTTTTGCCCACTTTCTGAATTAGGTGTAAGTAAAATTATACTCACTG 799
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This invention relates to novel i-antigen polypeptide sequences. I-antigens or immobilisation antigens ar common to a variety of hymenostomatid ciliates and their expression varies in response to
                                                                                                                                               PGKSQCTPCPAIKPANVAQATLGNDATITAQCNVACPDGTISAAGVNNWVAQNTECTNCA
                                                                                                                                                                                        846 CATAACACCAGAACAAGGGGTTG--CACACCCTAATTGGTCTATGGGTAAAAATTTCT
                                                                                                                                                                                                                                                                                                                                                                                                  371 TATLIA---QCALECPAGTVLTDGTTSTYKQAASECVKCAANFYTTKQTDWVAGIDTC--
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      48kD i-antigen repeat amino acid sequence SEQ ID 12.
                                                                                                        800 GTTCTGGCGGACCATTCCGTTACACGCCACTTGAACAATTCACCAA--
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Figure 5a; 144pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     UNIV GEORGIA RES FOUND INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ¥.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORNELL RES FOUND INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ichthyophthirius multifiliis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             99US-0118634.
99US-0122372.
99US-0124905.
99US-0131121.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAB25865 standard; Protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     04-FEB-2000; 2000WO-US02962
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1046 TCATTACTCAAATGGGA 1062
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       426 ----TSCNKKLTSGAEA 438
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Dickerson HW,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLARK T G.
DICKERSON H W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2000-506071/45.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           white spot disease;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     [mmobilisation
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02-MAR-1999;
17-MAR-1999;
27-APR-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAB25865;
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                                                                     168
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chthyophthirius multifilis, a protozoan which is an obligate parasite chthyophthirius multifilis, a protozoan which is an obligate parasite of freshwater fish causing ichthyophthiriasis or white spot disease. The convention includes two polypeptide and polynucleotide sequences for two i-antigens, of 48 and 55 kb. Also included in the invention are antibodies capable of binding to the nucleotide sequences and a method conferming the i-antigen nucleotide sequences. A composition (containing the i-antigen nucleotide) capable of eliciting an immune response in fish is useful for prophylaxis, treatment or for controlling I multifillis infection in fish. Polynucleotide or protein vaccines comprising a portion of the amplified product encoding an entigenic i-antigen polypeptide obtained is also useful for treating or preventing I multifillis infection in fish. Sequences AAA97036 A87042, and AAA97065 and AAA97080 represent i-antigen genes and gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to an isolated nucleic acid detection reagent capable of detecting 1000 or more genes from Drosophila. The invention is
                                                                                                                                                                                                                                                                                                                                                                                                                                          fragments identified in the invention. Sequences AAA97043-A97004 (excluding AAA97060) and AAA97071-A97088 represent primers used in the isolution of the i-antigen gene sequences. Sequences AAB25859-B25889 and AAB25893-B25906 represent i-antigen protein and peptide sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New isolated nucleic acid detection reagent for detecting 1000 or more genes from Drosophila and for elucidating cell signalling and cell-cell interactions -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         382 CPAGTVLIDGITSTYKQAASECVKCAANFYTTKQIDWVAGIDTCISCNKKLISGAEANLP 441
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Drosophila; developmental biology; cell signalling; insecticide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 72;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    8.3%; Score 212; DB 21;
53.6%; Pred. No. 1.4e-09;
tive 12; Mismatches 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Myers EW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ABB64261 standard; Protein; 3396 AA.
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11-JUL-2000; 2000US-0614150
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            37; Conservative
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N-PSDB; ABL08364.
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               442 ESAKKNIQC 450
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         61 AEATQKVQC 69
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   72 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 pharmaceutical.
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useful in developmental biology and in elucidating cell signalling and cell.cell interactions in higher eukaryotes for the development of insecticides, therapeutics and pharmaceutical drugs. The invention discloses genomic DNA sequences (ABL16176-ABL30511), expressed DNA (ABB57737-ABB12072).

The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences.
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1646 IDDLLNIANIGN-------QCPPLRALKSQISRGFNCNVGEVLNMDTSD 1687
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1731 TYTKEEGTKSQADCIPVCGYGTYSPTGL-----VPCLECPRNSF-----TAEPPTGG 1777
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1778 FKDCQACPAQSFTYQPA-----ASNKDLCRAKCAPGTYSATGL-----APCSP 1820
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1871 CVPMGHDIQCF-CPAG--FSGRRCEQDIDECASQPCYN------GGQCKDLPQGYRC- 1918
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1821 CPLHHYQGAA-----GAQSCNECPSNMRTDSPASKG-----REQCKPVVCGEGACQHGGL 1870
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1919 -ECPAGY-----SGINCQEEASDCGNDTCPARAMCKNEPGYKNVTCLCRSGYTGDQCD-- 1970
                                                                                                                                                                                                                                                                                                                                                                                                                         88 VIQCNVKCPAGTAIAGGATDYAAIIIECVNCRINFYNENAPNFNAGASTCTACPVNRVGG 147
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          148 ALTAGNAATIVAQCNVACPTGTALDDGVTTDYVRSFTECVKCRLNFYYNGNNGNTPFNPG 207
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            208 KSQCTPCPA----IKPANVAQATLGNDATITAQCNVACPDGTISAAGVNNWVAQNTECTN 263
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              264 CAPNEYNNNAPNFNPGNSTCLPCPANKDYGAEATAGGAATLAKQCN-1ACPDGTAIASGA 322
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              323 TNYVILQTECLNCAANFYFDGNNFQAGSSRCKACPANKVQGAVATAGGTATLIAQ---CA 379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                380 LECPAGTVLTDGTTSTYKQAASEC------VKC-AANFYTJKQTDWV 419
                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                         35 VDDLGTPANCVNCQKNFYINNAAAFVPGASTCTPCPQKK---DAGAQPNPPATANL---- 87
                                                                                                                                                                                                                                                       Ouery Match
8.1%; Score 206; DB 22; Length 3396;
Best Local Similarity 22.8%; Pred. No. 6.7e-07;
Matches 103; Conservative 37; Mismatches 181; Indels 130;
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